

Assay ID:

Sample name: Pyridoxine HCI Experiment start time: 9/12/2017 7:57:21 PM **UV-metric psKa** Analyst: Assay name: **Dorothy Levorse**

Instrument ID: 171-12014 T311053

Filename: C:\Sirius_T3\Mehtap\20170912_test_comp_pKa_diffexp\17I-12014_Pyridoxine HCI_UV-metric psKa.t3r

Yasuda-Shedlovsky result

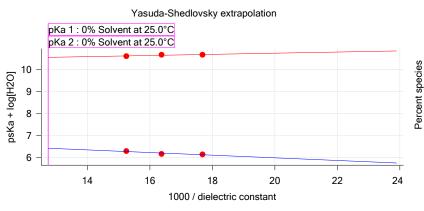
Extrapolation type pKa 0% SD Intercept Slope R^2 Ionic strength Temperature

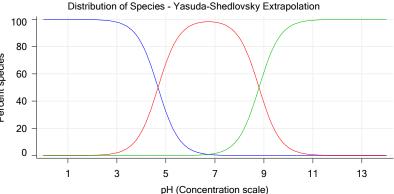
Yasuda-Shedlovsky 4.68 ±0.10 7.19 -60.2824 0.8298 0.165 M 25.0°C Yasuda-Shedlovsky 8.80 ±0.07 10.22 25.9224 0.6711 0.165 M 25.0°C

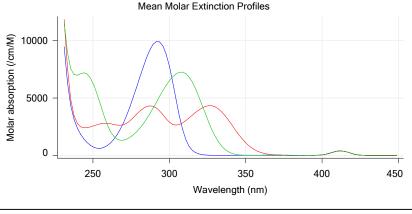
Component assay results

Titration	Methanol	Direction	Result	Dielectric	[H2O]	Ionic	Temperature		psKa	psKa
	weight%		type	constant		strength	-		1	2
17I-12014 Points 4 to 37	49.62 %	Up	UV-metric pKa	56.5	24.6 M	0.157 M	25.0°C	V	4.75 🔽	9.27
17I-12014 Points 39 to 79	39.80 %	Up	UV-metric pKa	61.1	30.1 M	0.166 M	25.0°C	V	4.68 🔽	9.19
17I-12014 Points 81 to 128	29.84 %	Up	UV-metric pKa	65.6	36.0 M	0.173 M	25.0°C	V	4.73 🔽	9.04

Graphs







UV-metric psKa Titration 1 of 3 17I-12014 Points 4 to 37

Results

pKa 1 4.75 pKa 2 9.27

RMSD 0.055 0.045 0.032

Chi squared 0.2284

PCA calculated number of pKas 2

Average ionic strength

0.157 M Average temperature 25.0°C

Analyte concentration range 30.5 μM to 28.8 μM

Methanol weight % Dielectric constant Water concentration 49.6 % 56.5

24.6 M



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Results (continued)

Number of pKas source Predicted

Wavelength clipping 230.0 nm to 450.0 nm pH clipping 1.469 to 12.530

Warnings and errors

Errors None

Warnings RMSD exceeds warning threshold

Assay Settings

Setting Value Original Value Date/Time changed Imported from

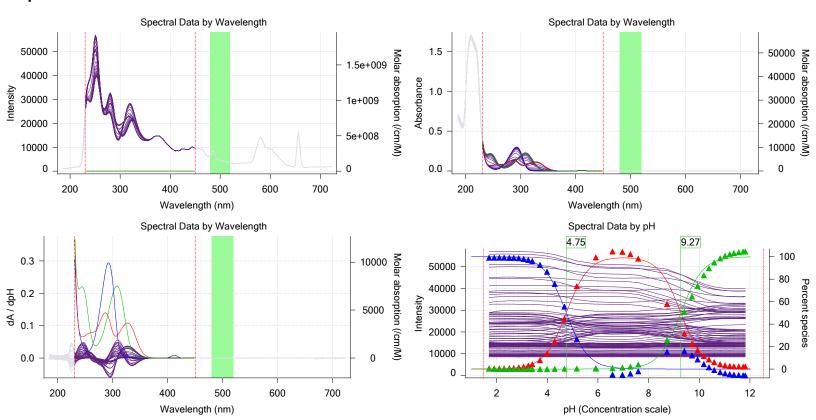
Buffer in use Ye Buffer type P

Phosphate Buffer

Assay Medium

Volume of buffer introduced 0.025000 mL Add buffer manually Manual

Graphs



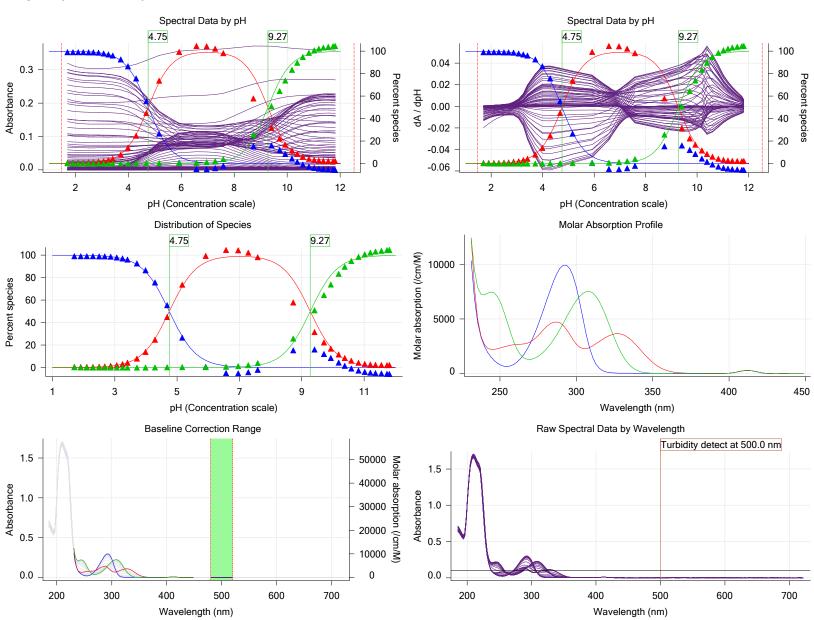


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Graphs (continued)



UV-metric psKa Titration 2 of 3 17I-12014 Points 39 to 79

Results

pKa 1 4.68 pKa 2 9.19 RMSD 0.084 0.071 0.049 Chi squared 0.3942

PCA calculated number of pKas 3

Average ionic strength
Average temperature
Analyte concentration range

0.166 M
25.0°C
25.0 µM

25.0 μM to 23.6 μM

Methanol weight % 39.8 % Dielectric constant 61.1 Water concentration 30.1 M



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Results (continued)

Number of pKas source Predicted

Wavelength clipping 230.0 nm to 450.0 nm pH clipping 1.445 to 12.525

Warnings and errors

Errors None

Warnings RMSD exceeds warning threshold

PCA calculation disagrees with predicted number of pKas

Assay Settings

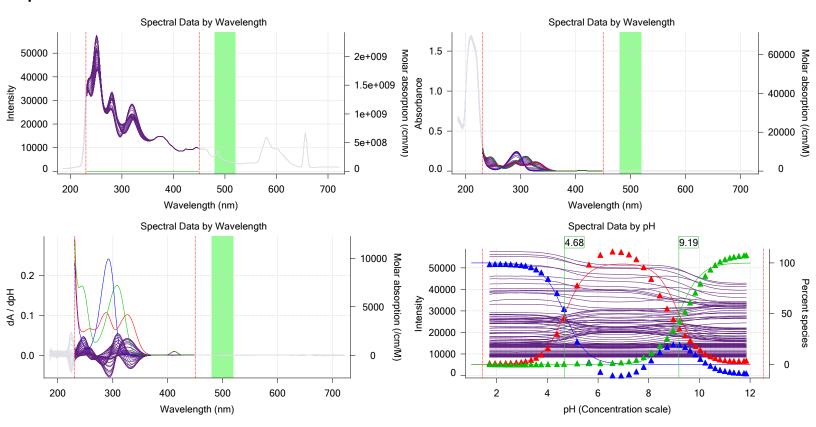
Setting Value Original Value Date/Time changed Imported from

Buffer in use Yes
Buffer type Phosphate Buffer

Assay Medium

Volume of buffer introduced 0.025000 mL Add buffer manually Manual

Graphs





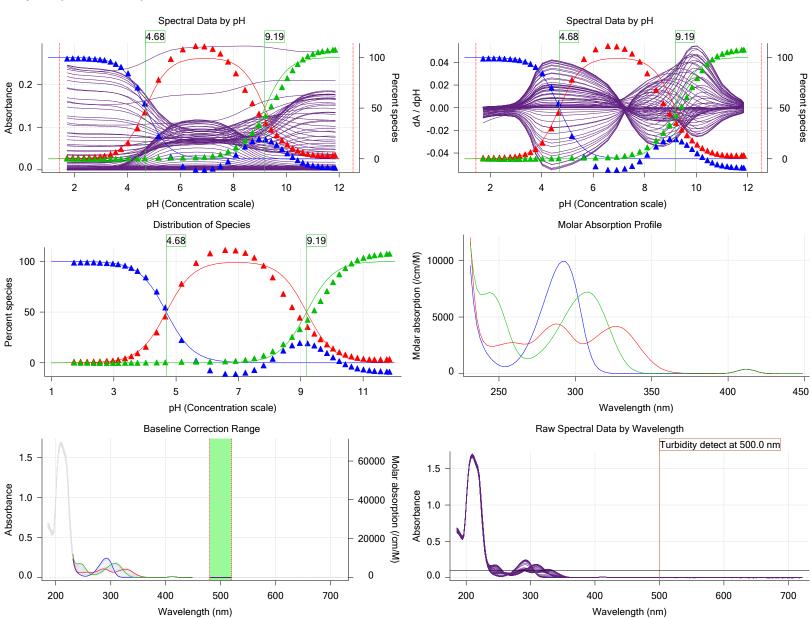
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Graphs (continued)



UV-metric psKa Titration 3 of 3 17I-12014 Points 81 to 128

Results

Water concentration

pKa 1 4.73 pKa 2 RMSD 0.088 0.066 0.061 Chi squared 0.5032 PCA calculated number of pKas Average ionic strength 0.173 M Average temperature 25.0°C Analyte concentration range 19.1 μM to 18.0 μM Methanol weight % 29.8 % Dielectric constant 65.6

36.0 M



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Results (continued)

Number of pKas source Predicted

Wavelength clipping 230.0 nm to 450.0 nm pH clipping 1.455 to 12.539

Warnings and errors

Errors None

Warnings RMSD exceeds warning threshold

PCA calculation disagrees with predicted number of pKas

Assay Settings

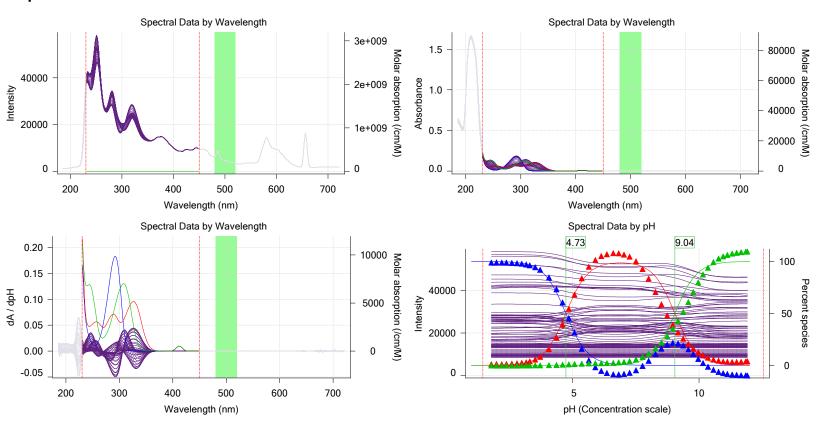
Setting Value Original Value Date/Time changed Imported from

Buffer in use Yes Phosphate Buffer Buffer type

Assay Medium

Volume of buffer introduced 0.025000 mL Add buffer manually Manual

Graphs



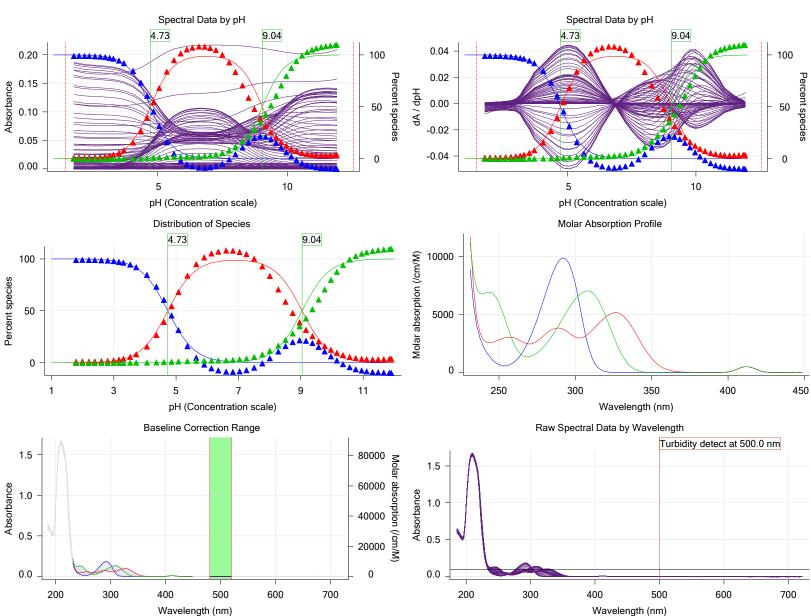


Experiment start time: 9/12/2017 7:57:21 PM Sample name: Pyridoxine HCI **UV-metric** psKa Assay name: **Dorothy Levorse** Assay ID:

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Graphs (continued)



Assay Model

nocay model			
Settings	Value	Date/Time changed	Imported from
Sample name	Pyridoxine HCI	9/12/2017 1:25:48 PM	User entered value
Sample by	Volume		Default value
Sample volume	0.0010 mL	9/12/2017 1:22:51 PM	User entered value
Solvent	DMSO		Default value
Sample concentration	0.048630 M	9/12/2017 1:22:51 PM	User entered value
Solubility	Unknown		Default value
Molecular weight	205.64	9/12/2017 1:23:01 PM	User entered value
Individual pKa ionic environments	No		Default value
Number of pKas	2	9/12/2017 1:22:51 PM	User entered value
Sample is a	Ampholyte	9/12/2017 1:22:51 PM	User entered value
pKa 1	4.90	9/12/2017 1:22:51 PM	User entered value
Туре	Base	9/12/2017 1:22:51 PM	User entered value
pKa 2	8.80	9/12/2017 1:22:51 PM	User entered value
Туре	Acid	9/12/2017 1:22:51 PM	User entered value



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Filename: $C:\Sirius_T3\\Mehtap\\20170912_test_comp_pKa_diffexp\\17I-12014_Pyridoxine\ HCI_UV-metric\ psKa.t3r$

Assay Model (continued)

Settings	vaiue	Date/Time changed	imported from
logp (XH2 +)	-10.00	•	Default value
logP (neutral XH)	-10.00	9/12/2017 1:22:51 PM	User entered value
logP (X -)	-10.00		Default value
Stoichiometry	1.00000		Default value
Aprotic counterion name	Chloride		From standards.xml file
Stoichiometry	1.00		From standards.xml file
Charge per counterion	-1		From standards.xml file

Stoichiometry Charge per counterion	1.00 -1		standards.xml file standards.xml file	
Assay Settings				
Setting	Value	Original Value	Date/Time changed	Imported from
General Settings				
Analyst name	Dorothy Levorse			
Separate reference vial	Yes			
Standard Experiment Settir	ngs			
Number of titrations	3			
Minimum pH	2.000			
Maximum pH	12.000			
pH step between points of	0.200			
Minimum titrant addition	0.00002 mL			
Maximum titrant addition	0.10000 mL			
Argon flow rate	100%			
Start titration using	Cautious pH adjust			
Advanced General Settings				
Detect turbidity using	Spectrometer			
Monitor at a wavelength of	500.0 nm			
Absorbance threshold of	0.100			
Collect turbidity sensor data	No			
Stir after titrant addition for	5 seconds			
For titrant addition, stir at	15%			
Titrant Pre-Dose				
Titrant pre-dose	None			
Assay Medium				
Cosolvent in use	Yes			
Cosolvent type	Methanol			
Cosolvent volume	1.15 mL			
Cosolvent added	Automatic			
ISA water volume	0.35 mL			
h				

After water addition, stir for At a speed of Buffer in use

Buffer type **Phosphate Buffer** 0.025000 mL Volume of buffer introduced

Automatic

5 seconds

15%

Yes

Add buffer manually Manual After medium addition, stir for 5 seconds

Sample Sonication Sonicate

Water added

No

Sample Dissolution

Perform a dissolution stage No

Carbonate purge

Perform a carbonate purge No

Temperature Control

Wait for temperature Yes Required start temperature 25.0°C Acceptable deviation 0.5°C Time to wait 60 seconds

Stir speed of 15%



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Assay Settings (continued)

Setting	Value	Original Value	Date/Time changed	Imported from
Titration 1				
Titrate from	Low to high pH			
Adjust to start pH	Yes			
After pH adjust stir for	10 seconds			
Titration 2				
Titrate from	Low to high pH			
Additional cosolvent volume	0.00 mL			
Add additional water	0.15 mL			
Additional water added	Automatic			
After pH adjust stir for	10 seconds			
Titration 3				
Titrate from	Low to high pH			
Additional cosolvent volume	0.00 mL			
Add additional water	0.34 mL			
Additional water added	Automatic			
After pH adjust stir for	10 seconds			
Data Point Stability				
Stir during data point collection	Yes			
For point collection, stir at	15%			
Delay before data point collection	0 seconds			
Number of points to average	20 points			
Time interval between points	0.50 seconds			
Required maximum standard deviation	0.00500 dpH/dt			
Stability timeout after	60 seconds			
Experiment cleanup				
Adjust pH to cleanup	To start pH			
And then stir for	60 seconds			
For cleaning, stir at	20%			
Then add water volume	0.25 mL			

Calibration Settings

And then stir for

Setting	Value	Date/Time changed	Imported from
Four-Plus alpha	0.080	9/12/2017 7:57:21 PM	C:\Sirius_T3\HCl17I12.t3r
Four-Plus S	1.0027	9/12/2017 7:57:21 PM	C:\Sirius_T3\HCl17l12.t3r
Four-Plus jH	1.2	9/12/2017 7:57:21 PM	C:\Sirius_T3\HCl17I12.t3r
Four-Plus jOH	-0.6	9/12/2017 7:57:21 PM	C:\Sirius_T3\HCl17l12.t3r
Base concentration factor	1.000		Default value
Acid concentration factor	1.000	9/12/2017 7:57:21 PM	C:\Sirius T3\HCl17l12.t3r

30 seconds

Instrument Settings

Setting	Value	Batch Id	Install date
Instrument owner	Merck		
Instrument ID	T311053		
Instrument type	T3 Simulator		
Software version	1.1.3.0		
Dispenser module		T3DM1100253	3/31/2009 6:24:52 AM
Dispenser 0	Water		3/31/2009 6:25:05 AM
Syringe volume	2.5 mL		
Firmware version	1.2.1(r2)		
Titrant	Water (0.15 M KCI)	8-18-17	9/8/2017 9:22:43 AM
Dispenser 2	Acid		3/31/2009 6:25:11 AM
Syringe volume	0.5 mL		
Firmware version	1.2.1(r2)		
Titrant	Acid (0.5 M HCI)	166940	9/8/2017 9:21:27 AM
Dispenser 1	Base		3/31/2009 6:25:21 AM



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Instrument Settings (continued)

Setting Syringe volume Firmware version	Value 0.5 mL 1.2.1(r2)	Batch Id	Install date
Titrant Dispenser 5 Syringe volume	Base (0.5 M KOH) Cosolvent 2.5 mL	01/06/17	9/8/2017 9:20:03 AM 3/31/2009 6:26:24 AM
Firmware version Distribution valve 5 Firmware version	1.2.1(r2) Distribution Valve 1.1.3		3/31/2009 6:28:19 AM
Port A Dispenser 3	Methanol (80%, 0.15 M KCl) Buffer	8-15-17	9/7/2017 3:40:52 PM 8/3/2010 6:05:16 AM
Syringe volume Firmware version Titrant	0.5 mL 1.2.1(r2) Phosphate Buffer		9/12/2017 12:32:29 PM
Dispenser 6 Syringe volume Firmware version	Octanol 0.5 mL 1.2.1(r2)		10/22/2010 11:52:43 AM
Titrant Titrator Horizontal axis firmware version	Octanol 1.17 Al1Dl2DO2 Stepper 2	T3TM1100153	9/12/2017 12:32:22 PM 3/31/2009 6:24:17 AM
Vertical axis firmware version Chassis I/O firmware version Probe I/O firmware version	1.17 Al1Dl2DO2 Stepper 2 1.11 Al1Dl0DO4 Norgren I/O 1.1.1		
Electrode E0 calibration	T3 Electrode -7.42 mV	T3E0769	8/15/2017 10:21:54 AM 9/12/2017 7:57:51 PM
Filling solution Liquids Wash 1	3M KCI 50% IPA:50% Water	KCL095	9/10/2017 10:38:19 AM 9/12/2017 9:11:25 AM
Wash 2 Buffer position 1 Buffer position 2 Storage position	0.5% Trition X-100 in H20 pH7 Wash pH 7		9/12/2017 9:11:30 AM 9/12/2017 9:11:33 AM 9/12/2017 9:11:37 AM 9/12/2017 9:11:42 AM
Wash water Waste Temperature controller Turbidity detector	8.6e+003 mL 1.5e+003 mL	9-11-17	9/11/2017 4:28:43 PM 9/11/2017 4:28:49 PM 8/5/2010 7:35:13 AM 3/31/2009 6:24:45 AM
Spectrometer Dip probe Wavelength coefficient A0	185.563	072390 11086	11/23/2010 12:22:28 PM
Wavelength coefficient A1 Wavelength coefficient A2 Total lamp lit time Calibrated on Integration time	2.17439 -0.000285622 75:06:25 9/6/2017 9:33:02 AM 11		11/23/2010 12:22:28 PM
Scans averaged Autoloader Left-right axis firmware version Front-back axis firmware version Vertical axis firmware version Chassis I/O firmware version	10 1.17 Al1Dl2DO2 Stepper 2 1.17 Al1Dl2DO2 Stepper 2 1.17 Al1Dl2DO2 Stepper 2 1.11 Al1Dl0DO4 Norgren I/O	T3AL1100237	11/10/2015 10:34:13 AM
Configuration Alternate titration position Alternate reference position Maximum standard vial volume Maximum alternate vial volume Automatic action idle period Titrant tube volume Syringe flush count Flowing wash pump volume	Titration position Reference position 3.50 mL 25.00 mL 5 minute(s) 1.3 mL 3.50 20.0 mL		



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Instrument Settings (continued)

Flowing wash stir duration Flowing wash stir speed Solvent wash stir duration Solvent wash stir speed Surfactant wash stir duration Surfactant wash stir speed E0 calibration minimum number of points E0 calibration maximum standard deviation E0 calibration timeout period E0 calibration stir duration E0 calibration preparation stir speed E0 calibration buffer wash stir duration E0 calibration buffer wash stir speed E0 calibration reading stir speed Spectrometer calibration stir speed Spectrometer calibration stir speed Spectrometer calibration wash pump volume Spectrometer calibration wash stir duration Spectrometer calibration wash stir speed	Value 5 s 30% 5 s 30% 5 s 30% 10 0.01500 60 s 5 s 30% 10000	Batch Id	Install date
Overhead dispense height	10000		

Refinement Settings

Setting	Value	Default value
Turbidity detection method	Spectrometer	Spectrometer
Turbidity wavelength to assess	500.0 nm	500.0 nm
Turbidity maximum absorbance	0.100	0.100
Turbidity probe threshold	50.00	50.00
Exclude turbid points	Yes	Yes
Low intensity warning threshold	100	100
Minimum absorbance change threshold	0.100	0.100
Eigenvector autocorrelation threshold	0.80	0.80
Maximum RMSD severe warning	0.250	0.250
Maximum RMSD warning	0.050	0.050